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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,768

DAIE: 03/27/2002

IIME: 13:24:50

Input Set A:\PK3065US3 SeqList.txt
Output Set: N:\CRF3\03272002\J083768.raw

SEQUENCE LISTING

		SEQUENCE LISTING
	4	(1) GENERAL INFORMATION.
	6	(i) APPLICANT: Dower, William J.
	7	Barrett, Ronald W.
	8	Cwirla, Steven E.
	Ğ	Duffin, David J.
	1(•	Gates, Christian
	11.	Haseiden, Sherril S.
	12	Mattheakis, Larry C.
	13	Schatz, Peter J.
	14	Wagstrom, Christopher R.
	15	Wrighton, Nicholas C.
	1 "	(ii) TITLE OF INVENTION: PEPTIDES AND COMPOUNDS THAT BIND TO A
	18	THROMBOPOIEIIN RECEPTOR
	20	(iii) NUMBER OF SEQUENCES: 232
	22	(iv) CORRESPONDENCE ADDRESS:
	23	(A) ADDRESSED: Glaxo Wellcome
	24	(B) STREET: Five Moore Drive, P.O. Box 13398
	25	(C) CITY: Research Triangle Park
	26	(D) STATE: NC
	2. '	(E) COUNTRY: USA
	28	(F) 2IP: 27709
	30	(V) COMPUTER READABLE FORM:
	3.1	(A) MEDIUM TYPE. Floppy disk
	32	(B) COMPUTER: LBM PC compatible
	33	(C) OPERATING SYSTEM. PC-DOS/MS-DOS
	3.4	(D) SOFTWARE: Patentin Release #1.0, Version #1.30
	30	(.1) CURRENT APPLICATION DATA:
C>		(A) APPLICATION NUMBER: US/10/083,768
C>	38	(B) FILING DATE: 27-Feb-2002
Ī	40	(viii) ATTORNEY/AGENT INFORMATION:
	41	(A) NAME: Hrubiec, Robert T
	4.2	(B) REGISTRATION NUMBER: 36 392
	43	10 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	4.5	(ix) TELECOMMUNICATION INFORMATION:
	45	(A) TELEPHONE: 919-248-1000
	49	
	5.1	(i) SEQUENCE CHARACTERISTICS:
	52	(A) LENGTH: 5 amino acids
	5.3	
	54	
	55	
	57	

DATE: 03/27/2002

TIME: 13:24:50

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,768

Input Set : A:\PK3065US3 SeqList.txt
Output Set: N:\CRF3\03272002\J083768.raw

```
(xi) SEQUENCE DESCRIPTION: SEQ ID No: 1:
W - - > 64
             Trp Ser Xaa Trp Ser
     6.5
             1
        (2) INFORMATION FOR SEQ ID NO: 2:
             (i) SEQUENCE CHARACIERISTICS:
     70
                  (A) LENGIH: 7 amino acids
     71
                  (B) TYPE: amino acid
                  (C) STRANDEDNESS:
     7 2
                  (D) TOPOLOGY linear
     35
            (:i) MOLECULE TYPE peptide
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
     80
W--> 82
             Xaa Xaa Xaa Xaa Xaa Xaa
     83
                              Ξ,
     85 (2) INFORMATION FOR SEQ ID NO: 3:
     87
             (i) SEQUENCE CHARACTERISTICS:
     88
                  (A) LENGTH: 9 amino acids
     89
                  (R) TYPE: amino acid
                  (C) STRANDEDNESS:
     90
     91
                  (D) TOPOLOGY: linear
     93
            (ii) MOLECULE TYPE: peptide
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
     98
              Xaa Gly Xaa Xaa Xaa Xaa Trp Xaa
W--> 100
     101
              1
     103 (2) INFORMATION FOR SEQ ID NO: 4:
              (i) SEQUENCE CHAFACTERISTICS
     16.5
     106
                   (A) LENGIH: 9 amino acids
     107
                   (B) TYPE: amano acad
     108
                   (C) STRANDELNESS:
     109
                   (D) TOPOLOGY, linear
             (ii) MOLECULE TYPE: peptide
     111
     116
             (xi) SEQUENCE DESCRIPTION. SEQ ID NO: 4:
W--> 118
              Xaa Gly Pro Thr Leu Arg Xaa Trp Xaa
     119
                              Ε.
     121 (2) INFORMATION FOR SEQ ID NO 5:
              (i) SEQUENCE CHAFACTEFISTICS:
     123
     124
                   (A) LENGIH: 10 am:no acids
     125
                   (B) TYPE: amono acid
     126
                   (C) STRANDEDNESS:
     127
                   (D) TOPOLOGY linear
     129
             (ii) MOLECULE TYPE, pertide
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
     134
W--> 136
              Xaa Xaa Gly Xaa Xaa Xaa Xaa Trp Xaa
                                                    10
     137
     139 (2) INFORMATION FOR SEQ ID NO 6:
     141
             (i) SEQUENCE CHARACTERISTICS
     142
                   (A) LENGTH: 18 amino acids
     143
                   (B) TYPE: amino acid
     144
                   (C) STRANDEDNESS:
     145
                   (D) TOPOLOGY. linear
```

RAW SEQUENCE LISTINGPATEL 03/27/2002 PATEL 03/27/2002 IME: 13:24:50

Input Set : A:\PK3065US3 SeqList.txt
Output Set : N:\CRF3\03272002\J083768.raw

```
(ii) MoLECULE TYPE peptide
147
        (d) SEQUENCE DESCRIPTION: SEQ ID Not 6:
152
         Gly Gly Cys Ala Asp Gly Pro Thr Leu Ard Glu Trp Tle Ser Phe Tys
1\rightarrow 4
                                               1.0
105
157
         Glv Gly
160 (2) INFORMATION FOR SEQ ID NO: 7:
        (i) SEQUENCE CHAFACTEFISTICS:
162
              (A) LENGIH 19 amino acids
163
              (B) TYPE: amino abid
1 4.4
              (C) STRANDEDNESS
165
1 41
              (D) TOPOLOGY: linear
        (:i) MOLECULE TYPE: pertide
168
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7
173
         Gly Asn Ala Asp Gly Pro Thr Leu Arg Gln Trp Leu Glu Gly Arg Arg
175
176
                                               10
         l
178
         Pro Lys Asn
191 (2) INFORMATION FOR SEQ ID NO: 8:
         (i) SEQUENCE CHARACTERISTICS:
              (A) LENGIH: 19 amino acids
184
              (B) TYPE: amino acid
185
186
              (C) STRANDEDNESS
              (D) TOPOLOGY, linear
187
189
        (ii) MOLECULE TYPE: pertide
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8
1:44
         Gly Gly Cys Aia Asp G.y Pro Ihr Leu Ard Glu Trp Ile Ser Phe Cys
146
                                               10
197
199
         Gly Gly Lys
202 (2) INFORMATION FOR SEQ ID NO: 9:
         (i) SEQUENCE CHAPACTERISTICS:
2014
2015
              (A) LENGIH: 18 amino acids
20h
              (B) TYPE: amino acid
202
              (C) STEANDEDNESS
2018
              (D) TOPOLOGY: linear
        (.i) MOLECULE TYPE: pertide
210
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 9
215
217
         Thr Ile Lys Gly Pro Thr Leu Arg Gln Trp Leu Lys Ser Arg Glu His
. 1 B
                                               10
220
         Thr Ser
223 (2) INFORMATION FOR SEQ ID NO: 10:
2 .....
         (i) SEQUENCE CHAFACIELISTICS:
              (A) LENGTH: 18 amino acids
226
237
              (B) TYPE: amino acid
              (C) SIFANDEDNESS:
22.8
              (D) TOPOLOGY linear
224
        (ii) MOLECULE TYPE: peptide
231
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO 10:
236
         Ser Ile Glu Gly Pro Thr Leu Arg Glu Trp Leu Thr Ser Arg Thr Pro
238
239
                                               10
         1
241
         His Ser
```

RAW SEQUENCE LISTING DATE: 03/27/2002 PATENT APPLICATION: US/10/083,768 TIME: 13:24:50

Input Set : A:\PK3065US3 SeqList.txt
Output Set: N:\CRF3\03272002\J083768.raw

```
244 (2) INFORMATION FOR SEQ ID NO. 11:
     246
              (i) SEQUENCE CHARACTERISTICS:
     _4<sup>-7</sup>
                    (A) LENGIH: 19 amino acids
     248
                    (B) TYPE amino acid
     249
                    (C) STFANDEDNESS:
     250
                    (D) TOPOLOGY linear
     252
              (ii) MOLECULE TYPE peptide
     257
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
              Leu Al: Ile Glu Gly Pro Thr Leu Arg Gln Trp Leu His Gly Asn Gly
     2.59
     200
              1
                               5
                                                    10
     252
              Arg Asp Thr
     265 (2) INFORMATION FOR SEQ ID NO: 12:
     267
              (i) SEQUENCE CHARACTERISTICS:
     글타꿈
                   (A) LENGTH: 14 amino acids
     263
                    (B) TYPE, amino acid
     27:)
                    (C) SIFANDEDNESS:
     200
                    (D) FOFOLOGY: linear
     273
             (1i) MOLECULE TYPE: peptide
     273
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
     230
              Cys Ala Asp Gly Pro Thr Leu Arg Glu Trp Ile Ser Phe Cys
     231
              1
                               ۲,
                                                    10
     283 (2) INFORMATION FOR SEQ ID NO: 13
     285
              (i) SEQUENCE CHARACTERISTICS
     286
                    (A) LENGTH: 14 amino acids
     187
                    (B) TYPE: amino acid
     288
                    (←) STFANDEDNESS:
     289
                    (D) TOPOLOGY: linear
     241
             (ii) MOLECULE TYPE, peptide
     246
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
     248
              lle Glu Gly Pro Thr Leu Arg Gln Irp Leu Ala Ala Arg Ala
     2.44
              1
                                                    10
     501 (2) INFORMATION FOR SEQ ID NO: 14
     303
              (i) SEQUENCE CHARACTERISTICS
     304
                    (A) LENGTH: 7 amino acids
     305
                    (B) TYPE: amino acid
     306
                    (C) STRANDEDNESS:
     3( "
                   (D) TOPOLOGY linear
     309
             (ii) MOLECULE TYPE: peptide
     314
             (xi) SEQUENCE DESCRIPTION SEQ ID NO: 14:
W--> 316
              Cys Xaa Xaa Xaa Xaa Xaa
     317
     319 (2) INFORMATION FOR SEQ ID NO. 15:
     321
              (i) SEQUENCE CHARACTERISTICS:
     322
                   (A) LENGTH: 16 amino acids
     323
                   (B) TYPE: amino acid
     324
                   (C) STRANDEDNESS:
     325
                   (D) TOPOLOGY: linear
     327
             (ii) MOLECULE IYPE, peptide
     332
             (xi) SEQUENCE DESCRIPTION. SEQ ID NO: 15:
```

RAW SEQUENCE LISTING DAIE: 03/27/2002 PAIENT APPLICATION: US/10/083,768 TIME: 13:24:50

Input Set : A:\PK3065US3 SeqList.txt
Output Set N:\CRF3\03272002\J083768.raw

```
Gly Gly Cys The Leu Ard 31t Trp Leu His Gly Gly Phe Cys Bly Gly
     4 3.1
                                                   . .
     4.45
     338 (2) INFORMATION FOR SEQ ID No: 16:
             (i) SEQUENCE CHARACTERISTICS:
     340
                   (A) LENGTH: 3 amino acids
     341
                   (B) IYPE: amino acid
     342
                   (C) SIRANDEDNESS:
     343
                   (D) TOPOLOGY: linear
     344
             (ii) MOLECULE IYPE: peptide
     34€
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
     351
              Xaa Cys Xaa Xaa Xaa Xaa Xaa
W - - > 353
                              5
     354
              1
     356 (2) INFORMATION FOR SEC ID NO 17:
              (i) SEQUENCE CHARACIERISTICS:
     358
                   (A) LENGIH: 15 amino acids
     359
                   (B) TYFE: amino acid
     360
                   (C) SIFANDEDNĒSS:
     361
                   (D) TOPOLOGY: linear
     362
             (ii) MOLECULE TYPE: peptide
     364
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
     364
              Ile Glu Gly Pro Thr Leu Arg Gln Trp Leu Ala Ala Arg Ala Lys
     372
                                                    1.0
     372
     3'4 (2) INFORMATION FOR SEQ ID NO: 18:
              (i) SEQUENCE CHARACTERISTICS:
     з"ь
     3 7 17
                   (A) IENGIH: 16 amino acids
                   (B) TYPE, arano acid
     378
     37+
                   (C) SIMANDEDNESS
                   (D) TOPOLOGY: linear
     38.4
             (ii) MOLECULE TYPE peptide
     38.
             (ix) FEATURE:
     335
                   (A) NAME, KEY Modified-site
     336
                    (B) LOCATION: 15
     337
                    (D) (THEF INFORMATION: /product= "Beta-ala"
     333
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
     391
              Ile Glu Gly Pro Thr Leu Arg Gln Trp Leu Ala Ala Arg Ala Xaa Lys
W - - > 393
                                                    10
                               ۳.
     391
              1
         (2) INFORMATION FOR SEC ID NO. 19
     33.
               (i) SEQUENCE CHARACTERISTICS.
     399
                    (A) LENGIH: 33 base pairs
     400
                    (B) TYPE, nucleic acid
     401
                    (3) SIRANDEDNESS, double
     402
                    (D) TOPOLOGY, linear
     403
              (ii) MOLECULE TYPE: GENA
     405
              (ix) FEATURE:
     403
                    (A) NAME KEY: CDS
     403
                    (B) LOCATION: 1..84
     410
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
     413
     415 CTC GAG AGC GGG CAG GTG GTG GAT GGG GAG CAG GTG GGT GAG GCC
                                                                                    48
     416 Leu Glu Ser Gly Gln Val Val His Gly Glu Gln Val Gly Glu Ala
```

VERIFICATION SUMMARY

PAIFNI APPLICATION: US/10/083,768

DAIE: 93/27/2002 TIME: 13:24:51

Input Set : A:\PK3065US3 SeqList.txt
Output Set: N:\CRF3\03272002\J083768.raw

```
1:37 M:226 C: Keyword misspelled or invalid format. [(A) APPLICATION NUMBER:]
1:38 M:220 C: Keyword misspelled or invalid format. ((B) FILING DAIE:)
L:64 M:341 W: +46 "n" or "Xaa" used, for SEQ ID=:1
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
1.1136 M 341 W _{\odot}(46) "n" or "Xaa" used, for SEQ ID= 5
L.316 M 341 W (46) "n" or "Xaa" used, for SEQ 1D#:14
                46 "n" or "Xaa" used, for SEQ ID# 16
1. 35: M :41 W
1:393 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#.18
L:45" M 541 W: (45) "n" or "Xaa" used, for SEQ ID#:21
L 460 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L 478 M·341 W (46) "n" or "Xaa" used, for SEQ ID#:22
L 481 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:22
L 499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 23
L 502 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:23
L:520 M:341 W: (46) "n" or "Xaa" used, for SEQ ID::24
L:523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L-1669 M-341 W: (40) "n" or "Xaa" used, for SEQ ID#.85
L:2812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:147
L. 2854 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#.149
I. 3322 M:341 W (4+) "n" or "Xaa" used, for SEQ IL# 172
                (40) "n" or "Xaa" used, for SEQ ID# 174
L-3362 M-341 W
                 (46) "n" or "Xaa" used, for SEQ ID# 175
L 3389 M:341 W.
L 3411 M.341 W (4) "n" or "Xaa" used, for SEQ ID# 176
L-3438 M-341 W- (46) "n" or "Xaa" used, for SEQ II #-17"
L 3465 M.341 W: (4) "n" or "Xaa" used, for SEQ II=178
I. 3487 M 341 W (4+) "n" or "Xaa" used, for SEQ ID#-179
L-3509 M-341 W (46) "n" or "Xaa" used, for SEQ ID#.180
L:3530 M:341 W (46) "n" or "Kaa" used, for SEQ ID#.184
L-3314 M-341 W (46) "n" or "Xaa" used, for SEO II # 196
I. 3854 M.341 W. (40) "n" or "Xaa" used, for SEQ ID# 198
I. 38)4 M 341 W. (46) "n" or "Xaa" used. for SE2 ID# 200
L.3934 M.341 W (4+) "n" or "Xaa" used, for SEQ ID# (202)
                (4) "n" or "Xaa" used, for SEQ \Pi(\#;2)4
I.: 3974 M. 341 W
                 i. "b" or "Yaa" used, for SED III# 223
L:1327 M 311 W
L:4345 M 341 W: (40) "n" or "Kaa" used, for SEQ ID# 224
L:4418 M 341 W: .460 "n" or "Kaa" used, for SE2 ID# 227
L:4433 M 341 W: (46) "n" or "Xaa" used, for SEQ ID= 228
L:4503 M 341 W: (46) "n" or "Xaa" used, for SEP ID#.231
L:4523 M.341 W: (4b) "n" or "Xaa" used, for SEQ ID::232
```